May 30, 2021

MAINSTREAM FINALLY CATCHES UP ON COVID-19 LAB LEAK THEORY - T.W, Z.L

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30th May 2021

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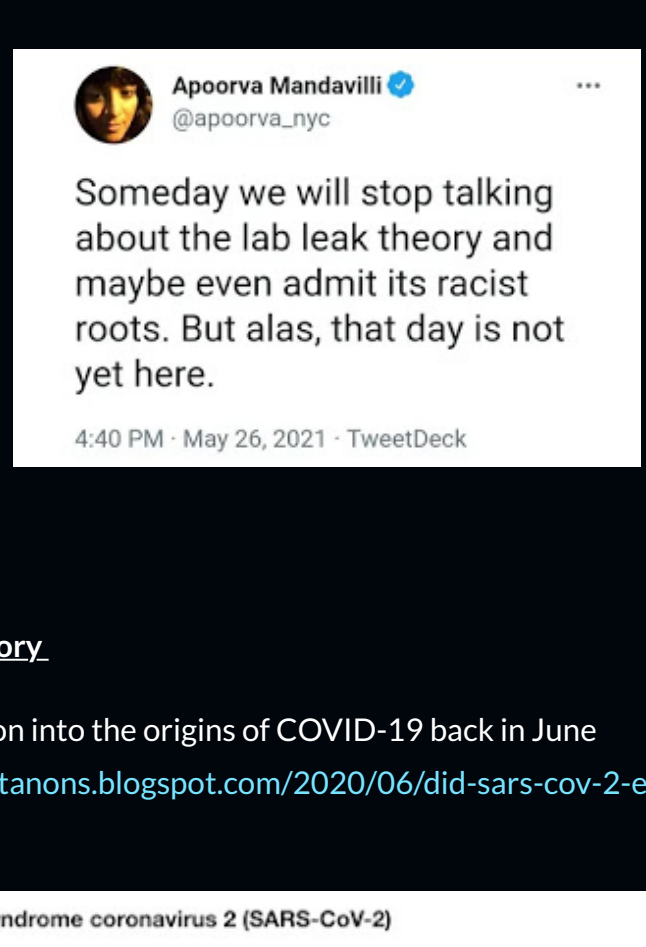


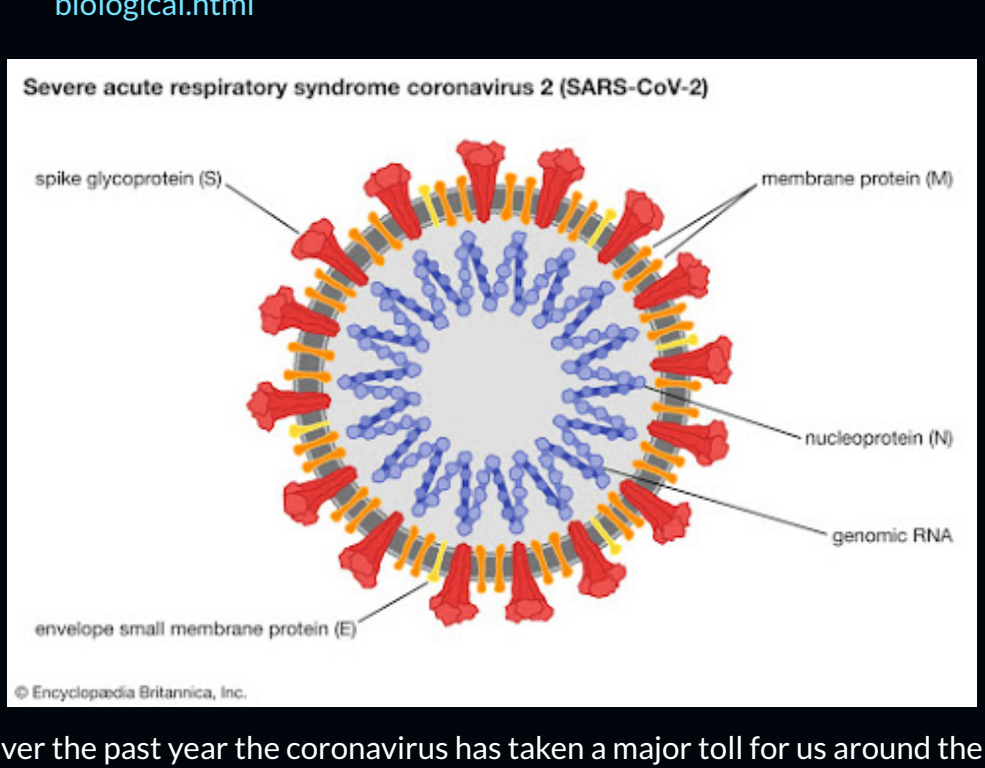
Apoorva Mandavilli
@apoorva_nyc

Someday we will stop talking about the lab leak theory and maybe even admit its racist roots. But alas, that day is not yet here.

4:40 PM · May 26, 2021 · TweetDeck

COVID-19 Lab Leak Theory

- See our Investigation into the origins of COVID-19 back in June 2020: https://cyclistanons.blogspot.com/2020/06/did-sars-cov-2-emerged-from-biological.html


Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)
spike glycoprotein (S)
membrane protein (M)
nucleoprotein (N)
genomic RNA
envelope small membrane protein (E)
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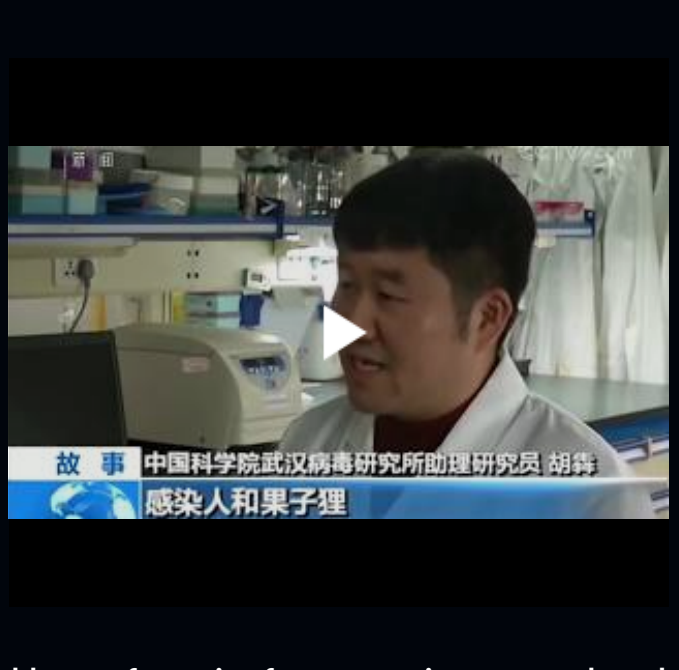
Over the past year the coronavirus has taken a major toll for us around the world, ranging from the millions of cases and deaths that may have involved friends and family, the new reality that has become a new normal and of course the trillions of dollars in economic damage plunged as a result of this. Most people didn't even induce the change until it was very late, and in most circumstances it wasn't easy to prepare against and adapt towards. For some time, the main conviction has been that the Severe Acute Respiratory Syndrome (abbreviated SARS-CoV-2 henceforth) which is universally known to cause COVID-19 disease, was that it occurred from a negligent spillover of hazardous bats that occurred within bat soup.

One year ago as of June 2020, we completed our independent investigation into the possible origin of COVID-19 that theorised that conceivably the outbreak of the disease originated in a manner contrary to the accepted narrative at the time. Most particular for our hypothesis, it was suggested that the virus may have been laboratory derived, introduced into Wuhan or may have already circulated for a longer time frame. For some time, the idea that SARS-CoV-2 was something outside zoonotic origins remained nothing more than a fringe conspiracy theory, however recently this has caught up to be more plausible in mainstream settings. The evidence for having natural zoonotic origins nonetheless continues to weigh in favour in contrast to the laboratory origin hypothesis, previously being a very minimally considered source of investigation in the scientific community.

To reiterate though, currently there is no conclusive evidence that SARS-CoV-2 was a natural zoonosis or even being artificially manufactured, therefore we can only speculate and outweigh the evidence on both sides to determine two equally feasible options. It has been deemed to have originated from a bat due to the genetic similarities it has towards bats, though it lacks the signs expected for natural emergence from the environment.

In May 2020, the World Health Assembly that governs WHO voted on a motion to pursue investigations into the origins of SARS-CoV-2, which received support from 137 member-states including China.
Based on the World Health Organisation's own investigation in conjunction with China, they have theorised firmly on the position that the virus occurred through a human spillover by a reservoir species.

Most of our past findings were heavily censored by Big Tech as part of their campaign to combat "misinformation", though we able to confirm a few more things that have built on its consensus. Let's look at some of the few rejected notions we arguably got correct though note this is still pending an investigation:



- **Given the arguably unsafe setting for coronavirus research at the Wuhan Institute of Virology, this increases the possibility of a lab leak being possible.**

It was concluded that both the Wuhan Institute of Virology (WIV) and Wuhan Centre for Disease Control and Presentation located in the city were already engaging in routine studies of coronaviruses including those of bat origin. Research done at the WIV were conducted using laboratories of Biosafety Level-2 and 3, it did however recently open up China's first Biosafety Level 4 laboratory being the safest. After the emergence of the initial COVID-19 outbreak, internal policies would change practices to limit the cultivation and experimentation of highly dangerous viruses within a BSL-4 laboratory.

Dr Peter Daszak, President of EcoHealth Alliance who has also openly endorsed his past work done in conjunction with the WIV, also has detailed the work conducted included the artificial programming of coronaviruses onto humanised mice. This is able noteworthy as humanised mice, which being engrafted with such human tissues and genes means it makes it possible for pathogens to transmit to humans (as experimental viruses are now programmed to target human cells, though in this case a lab rat bearing human substances is substituted).

According to WIV virologist Shi Zhengli, it was claimed there were no reported incidents of infection among any staff or students when handling bats. Contrary to this assumption, it was reported by US Intelligence that there were at least three researchers that were hospitalised around Chinese Autumn season of 2019, exhibiting symptom signs consistent with the coronavirus disease. The first public notice provided to the international community regarding the coronavirus wouldn't occur until December 31st 2019 when then World Health Organization office in China was notified.

In terms of bats transmitting the virus, in most cases it can occur through bat saliva transmission either through a direct bite, scratch or any mucous membrane exposure. It is also noted that there has been history of its researchers being bitten by bats, which upon observation of working conditions shreds concern for accidentally unwitting exposure. Despite this and the fact that it was reported that no specific incidents of infection from bats, the images previously hosted on the WIV' website over the time before the initial outbreak highlights concern for lack of reinforcing protective measures despite past instances. Notice the lack of personal protective equipment (PPE) including the image on the left of bats being handled without biohazard gloves:

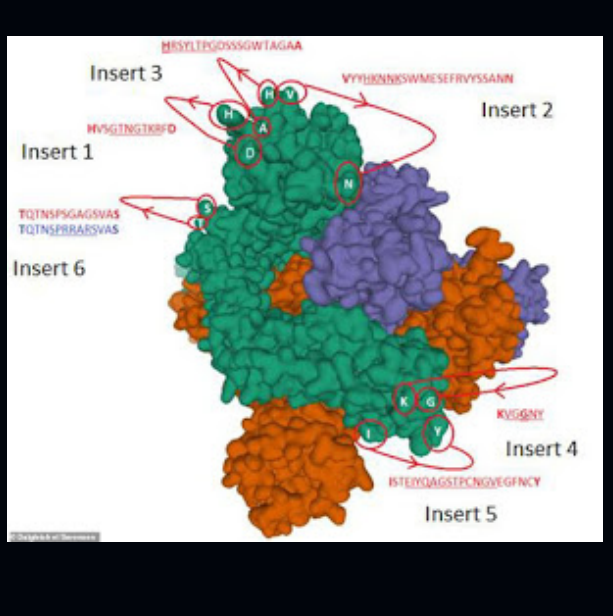
Article: 【中国青年报】"蝙蝠女侠"团队找出SARS病毒源
http://www.whiov.cas.cn/xwdt_105286/kydt/201712/t20171214_4916301.html

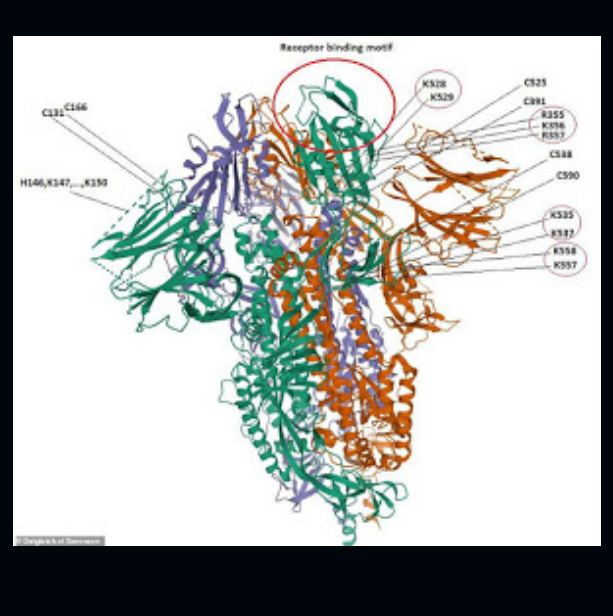
Similar to bees, bats tend to refrain from malicious contact with humans unless directly provoked, which can result in aggression. Medical advise furthermore strictly advises even against touching bats with bare hands due to the prospect of coming into contact with infectious material. The prospect of an infected bat attacking researchers even within the controlled environment of a laboratory does remain possible, which an out of control bat spews into chaos and during the struggle to restrain the creature, spills something infectious.

Based on a report by researchers in Wuhan Centre for Disease Control and Prevention (just across the road from the suspected seafood market), there has been cases where a researcher has experienced attacks from bats, prompting the need to volunatrily quarantine themselves due to the risk. Through a JAMA network study as early as February 2020, it has been determined that recovered patients can still possess the risk of being asymptomatic even AFTER symptoms cease to exhibit themselves. It is also confirmed that on November 18th 2019, a job posting was also published to recruit postgraduate students, which would have involved the study of the molecular mechanisms associated with coronaviruses continuing to host itself on asymptomatic bodies.

- **SARS-CoV-2 having origins of an artificially manipulated nature**

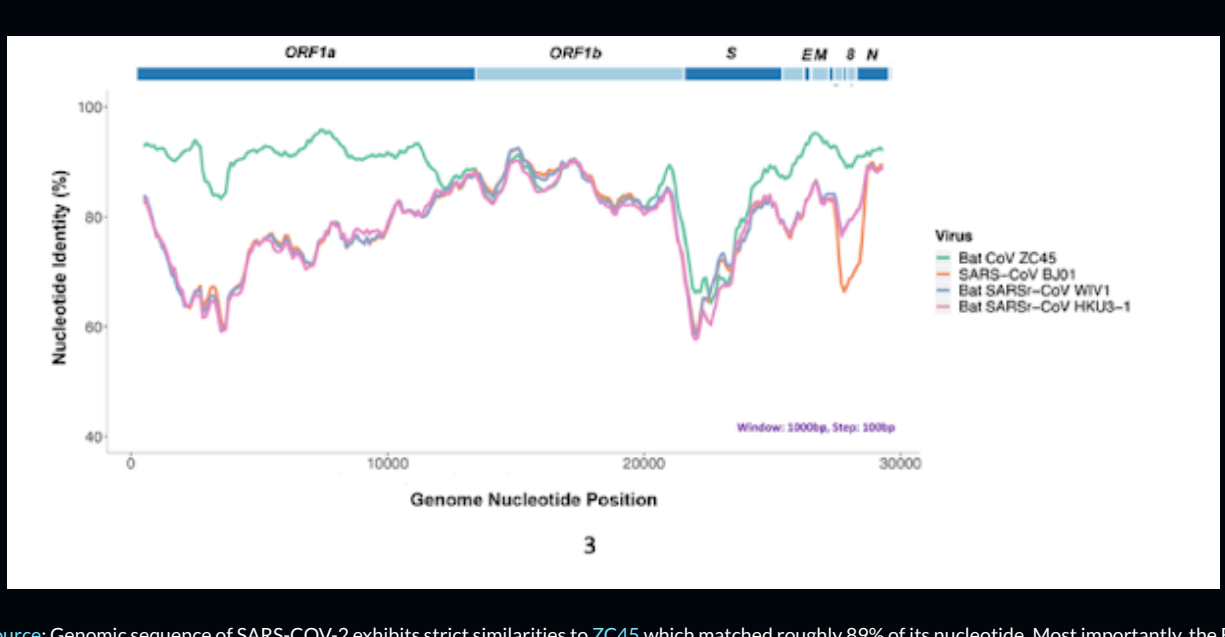
For a significant portion of the pandemic and which has continued to be recognised by the World Health Organization, it was accepted that the origins of COVID-19 derived from bats. As previously mentioned in our past investigation, horseshoe bats located in Wuhan, China remains arguably a difficult process. This is particularly due to the urban environment that generally deters bats and the fact that during the temperatures were low at the time in Wuhan, bats naturally would have been expected to be hibernating. Furthermore, Wuhan is also located relatively distant from natural settings where bats typically are present, meaning any bats would have required importation, begging the question on possible cases outside the city beforehand. Some past testimonies have also suggested that the Huanan Seafood Wholesale Market also were selling little to no bats at the time of the outbreak. For the Wuhan Institute of Virology, most bats were retrieved from a cave in Yunnan, China that were sampled for coronaviruses and used for lab experiments.





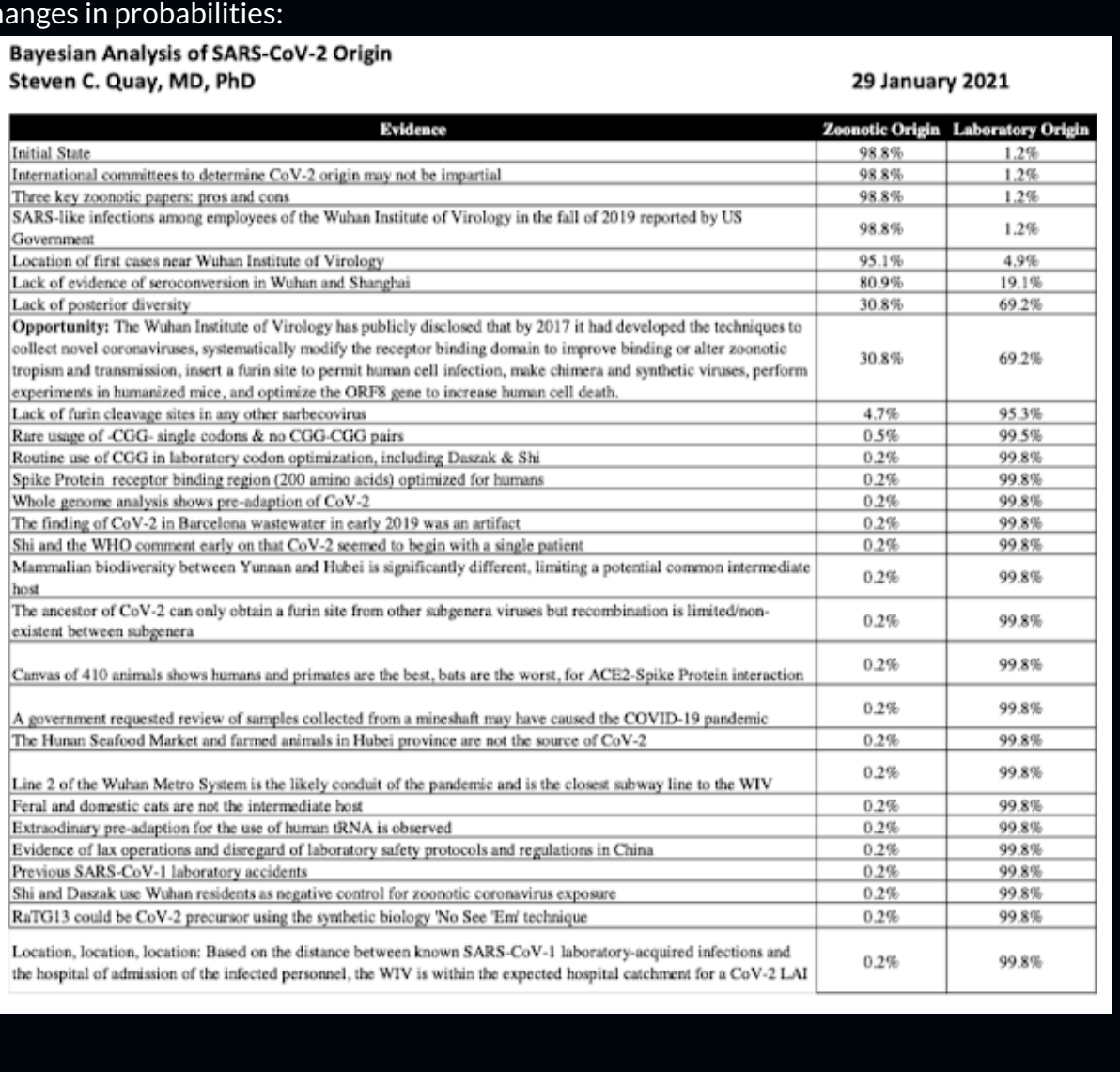
(Left: Four changing genomic 'fingerprints' suggesting the ordering of the DNA and RNA of the coronavirus that would possibly require laboratory manipulation with that precision, Right: Four amino acids found on the SARS-Cov-2 spike protein have a positive charge that can transfer to human cells)

It has also been shown that roughly around 70-96% of the genetic sequences associated with the 2019-novel coronavirus bears a high degree of similarity with bat viruses possessed at the Wuhan Institute of Virology.



(Source: Genomic sequence of SARS-COV-2 exhibits strict similarities to ZC45 which matched roughly 89% of its nucleotide. Most importantly, the E-Protein was 100% identical confirming it is homologous, it also showed that the Nucleocapsid protein was 94% identical, the Membrane protein is 98.6% identical, the second half of the Spike protein was 95% identical and the ORF8 gene was 94.2%. Biggest difference was SARS-COV-2 was it contained receptor-binding motif in it's spike protein, which could bind hACE2 which Angiotensin-converting enzyme 2 in the human organs could allow entry and digestion of infection. The original ZC45 was a research on bats back in 2015-17 at Chinese military laboratories in Chongqing and Nanjing, which over two decades there has been a collection of bat coronavirus backbone templates)

The following Text-Table from within a Bayesian Analysis summarises the evidence examined and the changes in probabilities:



Bayesian Analysis of SARS-CoV-2 Origin
Steven C. Quay, MD, PhD
29 January 2021

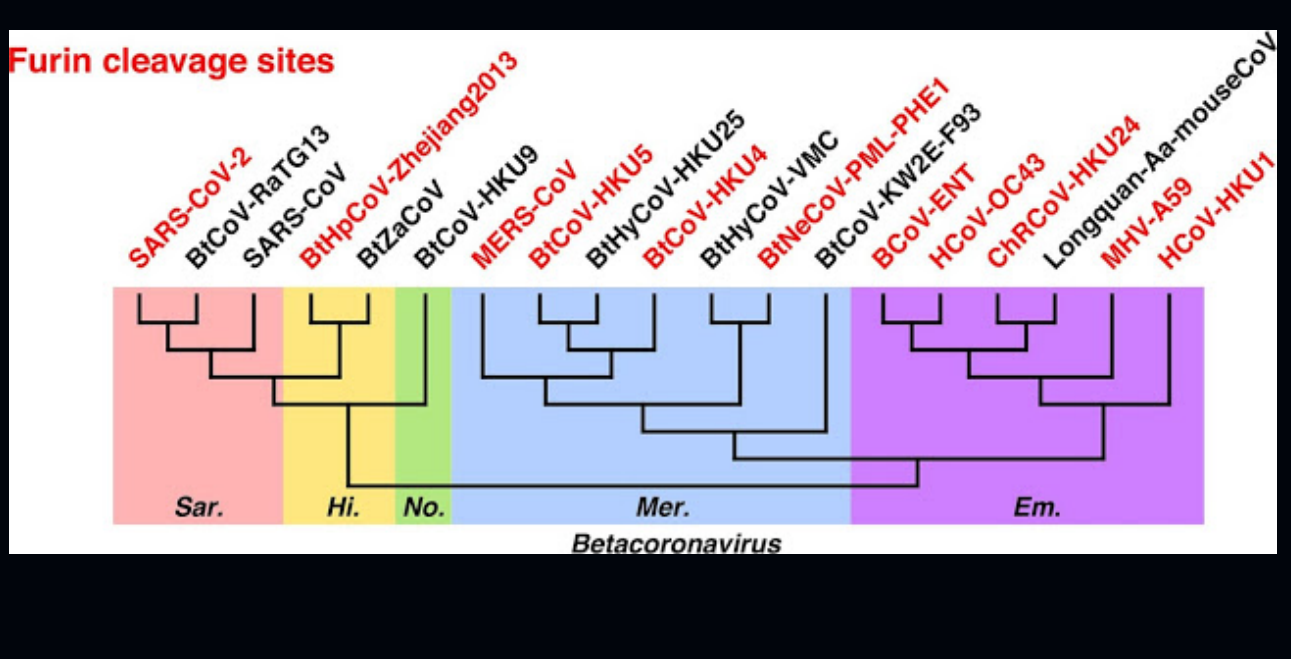
Evidence	Zoonotic Origin	Laboratory Origin
Initial State	98.8%	1.2%
International committees to determine CoV-2 origin may not be impartial	98.8%	1.2%
Three key zoonotic papers: pros and cons	98.8%	1.2%
SARS-like infections among employees of the Wuhan Institute of Virology in the fall of 2019 reported by US Government	98.8%	1.2%
Location of first cases near Wuhan Institute of Virology	95.1%	4.9%
Lack of evidence of seroconversion in Wuhan and Shanghai	80.9%	19.1%
Lack of posterior diversity	30.8%	69.2%
Opportunity: The Wuhan Institute of Virology has publicly disclosed that by 2017 it had developed the techniques to collect novel coronaviruses, systematically modify the receptor binding domain to improve binding or alter zoonotic tropism and transmission, insert a furin site to permit human cell infection, make chimera and synthetic viruses, perform experiments in humanized mice, and optimize the ORF8 gene to increase human cell death.	30.8%	69.2%
Lack of furin cleavage sites in any other sarbecovirus	4.7%	95.3%
Rare usage of -CGG- single codons & no CGG-CGG pairs	0.5%	99.5%
Routine use of CGG in laboratory codon optimization, including Daszak & Shi	0.2%	99.8%
Spike Protein receptor binding region (200 amino acids) optimized for humans	0.2%	99.8%
Whole genome analysis shows pre-adaption of CoV-2	0.2%	99.8%
The finding of CoV-2 in Barcelona wastewater in early 2019 was an artifact	0.2%	99.8%
Shi and the WHO comment early on that CoV-2 seemed to begin with a single patient	0.2%	99.8%
Mammalian biodiversity between Yunnan and Hubei is significantly different, limiting a potential common intermediate host	0.2%	99.8%
The ancestor of CoV-2 can only obtain a furin site from other subgenera viruses but recombination is limited/non-existent between subgenera	0.2%	99.8%
Canvas of 410 animals shows humans and primates are the best, bats are the worst, for ACE2-Spike Protein interaction	0.2%	99.8%
A government requested review of samples collected from a mineshaft may have caused the COVID-19 pandemic	0.2%	99.8%
The Hunan Seafood Market and farmed animals in Hubei province are not the source of CoV-2	0.2%	99.8%
Line 2 of the Wuhan Metro System is the likely conduit of the pandemic and is the closest subway line to the WIV	0.2%	99.8%
Feral and domestic cats are not the intermediate host	0.2%	99.8%
Extraordinary pre-adaption for the use of human tRNA is observed	0.2%	99.8%
Evidence of lax operations and disregard of laboratory safety protocols and regulations in China	0.2%	99.8%
Previous SARS-CoV-1 laboratory accidents	0.2%	99.8%
Shi and Daszak use Wuhan residents as negative control for zoonotic coronavirus exposure	0.2%	99.8%
RaTG13 could be CoV-2 precursor using the synthetic biology 'No See 'Em' technique	0.2%	99.8%
Location, location, location: Based on the distance between known SARS-CoV-1 laboratory-acquired infections and the hospital of admission of the infected personnel, the WIV is within the expected hospital catchment for a CoV-2 LAI	0.2%	99.8%

What makes SARS-CoV-2 unique is the lack of creditable ancestry associated with the virus, most particularly it appears intensively pre-adapted to human cells in contrast to previous viral outbreaks of a similar nature. Most particularly, SARS-CoV-2 was reported as of May 2020 to have the advantage within a position equivalent to the late SARS-CoV epidemic where it had already evolved towards a lethal human pathogen. Specifically its spike protein had already reached the stage where 99.5% the amino acid sequence could already be applied to the human receptor. At a genomic level, three other Orthornavirae coronaviruses would be identified as matching SARS-CoV-2, including Pangolin-CoVs (2017, 2019), Bat-SARS-like (CoVZC45, CoVZXC21) and bat RatG13. This helps reinforce the zoonotic origin claim, given the sequence of SARS-CoV-2 matches that of bats and pangolin.

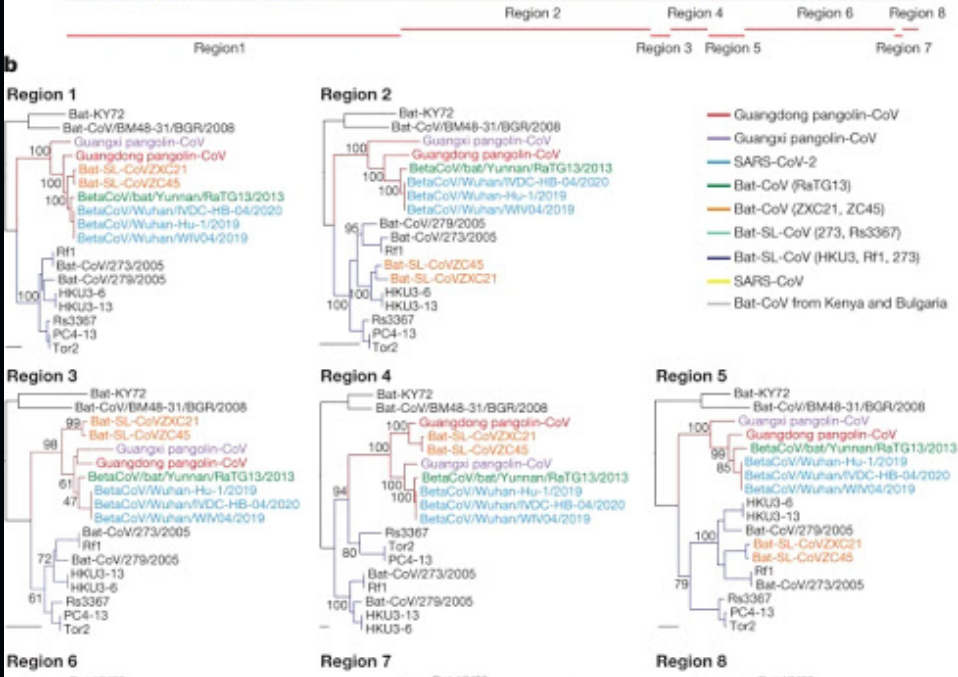
What makes SARS-CoV-2 unique is that the novel furin cleavage site in the spike glycoprotein that increases the infectivity from the virus anatomy, at this site is also facilitated by a polyphasic insertion that the furin can cleve. This helps flank the suboptimal furin CS of SARS-CoV-2 spike to enhance the viral transmission into the mucosal epithelial and primary human airway cells. The Proteolytic cleavage associated with the spike glycoprotein would also help determine on whether the virus is able to bind onto human cells and particularly mediate entry.

Encoded within the spike glycoprotein there is 12 nucleotides inserted over four amino acids PRRA over the furin cleavage site. For this insertion to occur, it would require a recombination by a donor from another furin site and virus, it does shred the possibility of laboratory insertion especially due to its rarity. Differentiating all this, we see CGG-CGG codons encoding the two arginines of the doublet within SARS-CoV-2 appear to be not visible in any other furin sites of the viral proteins expressed by a range of similar subgenera viruses. Coronaviruses in nature if occurred naturally would make this extremely rare, as the pairing of CGG-CGGG codons has been known only to have a <1% probability.

Even the highly related bat-virus CoV RaTG-13 does not appear to have featured a furin cleavage site. Ultimately what is been seen appears to have very unique characteristics exclusive to SARS-CoV-2 given the cleavage site would have required such recombination, naturally or artificially. In the virology community, it is also typically known that if a virus was to be precipitated to a more lethal condition then it could involve placing a furin cleavage site at the S1/S2 subunits of the junction, both being key domains for receptor binding and enabling membrane fusion.



Changing Patterns and Sequence Similarities Between SARS-Cov-2 and Bats/Pangolins Coronaviruses



- **Gain-of-Function Coronavirus Research**

Back in October 2014, the US Government under the Obama Administration issued a federal moratorium against gain-of-function research, in response to concerns regarding the country's biosafety measures and standards. This followed after series of laboratory accidents at the Centre for Disease Control in July of that year, including exposure to viruses and the contamination of samples. The response predominately came with over 200 scientists signing the Cambridge Working Group Declaration to petition the temporarily cession on experiments involving the manufacturing of viral pathogens. As part of a public health measure, this suspended all public funding for any research relating to the biological engineering of weaponising viruses.

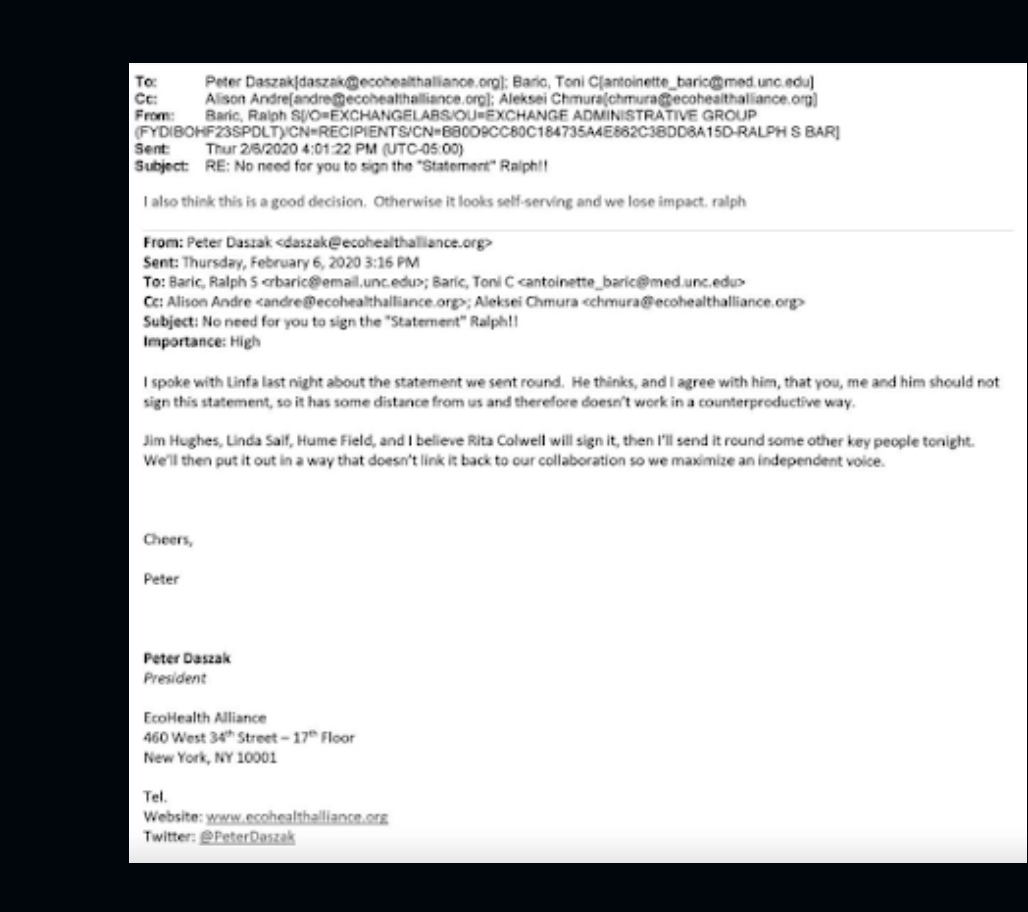
The following year in 2015, an alternative approach was taken in response to the restrictions imposed within the United States by the National Institute of Allergy and Infectious Diseases Director Anthony Fauci. Gain-of-function research was then outsourced to China's Wuhan Institute of Virology to license it's laboratory, authorising a US Government grant to be utilised by nongovernmental research group EcoHealth Alliance. These grants to the EcoHealth Alliance were awarded by the NIH and NIAID, with the prospect that a project would be run in collaboration within Wuhan for engineering viruses.

- Understanding the Risk of Bat Coronavirus Emergence Grant

These funds would be designated to Wuhan Institute for the purposes of collecting and analysing viral samples. It was expected that by genetically engineering a virus, this would enable researchers to concurrently observe and further commence any development of vaccines that could cure those viruses. Up to $3.7 million was granted over the duration of the project, and was later renewed in 2019 before eventually having another grant approved but then terminated by the Trump Administration by May 2020.

The National Institute of Allergy and Infectious Diseases eventually advised the White House to lift the ban in 2017. This came without any notification towards senior White House officials of the risks of accidents sparking a pandemic, which at the time Fauci reaffirmed that it was "important work" to outweigh the risk. In the past, Dr Fauci has defended the prospect of gain-of-function research despite the high risk of a man-made pandemic. Basically justifying that on the premise that an accidental pandemic could occur, the benefits gained from the experimentation will have greater benefit. Most proponents of this argued that gain-of-research function's role in facilitating vaccine development, would ensure optimal and timely preparation of such a vaccine should any public health disaster occur.

As EcoHealth Alliance was active in Wuhan with their research even upon the emergence of a suspected pneumonia outbreak, a very pressing statement was published in February 2020 to publicly inform that "We stand together to strongly condemn conspiracy theories suggesting that COVID-19 does not have a natural origin". The thing was that it was still early on the outbreak, in addition on the idea that laboratory leaks do frequently happen it was immediately rejected for any further observation, and that it was published by EcoHealth Alliance President Peter Daszak. Through an email uncovered, we see that there was efforts pushed to conceal any potential conflict of interest that might be associated with EcoHealth Alliance. We see this as part of the group's effort to deflect any responsibility or visible connection, or which Daszak would later refer as "crackpot theories that need to be addressed" in relation to theories of laboratory incident.



To: Peter Daszak[daszak@ecohealthalliance.org]; Baric, Toni C[antoinette_baric@med.unc.edu]
Cc: Alison Andre[andre@ecohealthalliance.org]; Aleksei Chmura[chmura@ecohealthalliance.org]
From: Baric, Ralph S[/O=EXCHANGELABS/OU=EXCHANGE ADMINISTRATIVE GROUP (FYDIBOHF23SPDLT)/CN=RECIPIENTS/CN=B80D9CC80C184735A4E862C3BDD8A15D-RALPH S BAR]
Sent: Thur 2/6/2020 4:01:22 PM (UTC-05:00)
Subject: RE: No need for you to sign the "Statement" Ralph!!

I also think this is a good decision. Otherwise it looks self-serving and we lose impact. ralph

From: Peter Daszak <daszak@ecohealthalliance.org>
Sent: Thursday, February 6, 2020 3:16 PM
To: Baric, Ralph S <rbaric@email.unc.edu>; Baric, Toni C <antoinette_baric@med.unc.edu>
Cc: Alison Andre <andre@ecohealthalliance.org>; Aleksei Chmura <chmura@ecohealthalliance.org>
Subject: No need for you to sign the "Statement" Ralph!!
Importance: High

I spoke with Linfa last night about the statement we sent round. He thinks, and I agree with him, that you, me and him should not sign this statement, so it has some distance from us and therefore doesn't work in a counterproductive way.

Jim Hughes, Linda Saif, Hume Field, and I believe Rita Colwell will sign it, then I'll send it round some other key people tonight. We'll then put it out in a way that doesn't link it back to our collaboration so we maximize an independent voice.

Cheers,

Peter

Peter Daszak
President

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New York, NY 10001

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Website: www.ecohealthalliance.org
Twitter: @PeterDaszak

What the Lab Leak Is Proven Beyond Reasonable Doubt?

In the slim chance that the lab leak hypothesis is irrefutably proven, how the international community would react would likely be based on whether there was any malicious intention to release a biological agent or if it had been an accidental circumstance (most leaks from laboratories are a common occurrence and concern). Even today, most unplanned economic or environmental damage that might indirectly impact another sovereign state is often dismissed, though it is likely other states rather then just China will face condemnation and pressure to reform specific public policies regarding laboratory research.

Since 1975, a multilateral disarmament agreement was signed by multiple countries which also included China as a signatory within the Convention on the Prohibition of the Development, Production and Stockpiling of Bacteriological (Biological) and Toxin Weapons and on their Destruction (Biological Weapons Convention). How this international treaty operates is through member-states to the treaty being able to submit complaints of BWC violations to the UN Security Council which would instigate an investigation against the accused member state. Limitations on what member-states are able to perform in regards to biological research with toleration by the international community relies on the expected standard. Even if China had engaged in bio-warfare research, then this would technically still be permissible as like any signature state are allowed to engineer biological weaponry within limitations to purposes such as defence.

Opinion) Proposed Theory on Origin:

- Past public health disasters have prompted research into SARS, which for China prompted it's plans to construct between 5-7 biosafety level 4 laboratories across mainland China by 2025. One of these prominent facilities that also included the Wuhan Institute of Virology that was approved by the Chinese Academy of Sciences in 2003, which would eventually become a hotspot to secure some of the world's most deadliest and largest pathogens. Pressure from the international community enables an injection of additional foreign aid to support China's initiative both for domestic and foreign usage.
- Investigations into the origins of SARS by researchers at the Wuhan Institute of Virology traced back to a horseshoe bat cave in Yunnan Province, China. Thousands of samples would be collected between 2005-2019 to be examined in Wuhan (over 1500+ kilometres away from the cave), which revealed several spike proteins capable of binding towards human ACE2.
- The RaTG13 virus is known to have originated from a copper mine in Moijang (also in Yunnan Province), specifically in 2012 when three miners working in the bat-infested location contracted bat-coronavirus. It was exhibited in a manner similar to COVID-19 symptoms and later the three were pronounced deceased. Samples were eventually brought to the Wuhan Institute of Virology where they would remain in storage for further testing up until 2020.
- Several gain-of-function research would eventually be contracted at Wuhan Institute of Virology both by the Chinese People's Liberation Army and other foreign actors through the lease of the facility. Despite presenting itself as a civilian research institution, China would strategically divert this funding including from donors to support military projects. Given the level of international oversight over the facility, China would conceal the true intent of its operations occurring at WIV, whilst commencing with classified research and experimentation. While the proliferation of manufactured viruses would also be weaponised, a biological weapon of such would be expected to remain on standby in a substituting manner like nuclear weapons.
- Many who had been working at the Wuhan Institute of Virology would accidentally leak one of the lethal virus samples due inadequate handling within the laboratory setting. One of the first individuals (likely a researcher) that contracted the virus would visit the Huanan Seafood Wholesale Market, where two-thirds of the early cases were exposed towards. It is also possible that one of the traders at the wet market recognised nearby biohazard waste (from the nearby BSL-4 labs) that included bat meat. This possibly without any regard and/or desperation may have caused the hypothetical individual to source and eventually distribute the bat to several patrons at Huanan Seafood Wholesale Market.
- At some stage in 2019, public health authorities in Wuhan discovered patient zero who had contracted the novel coronavirus SARS-CoV-2. Efforts would be made to censor any records and public accounting relating to this suspected patient, including shielding it's presence from the Central Committee of the Chinese Communist Party in Beijing, China. Aggressive efforts would be made to silence any possible whistleblowers, including upon discovery that the virus had genetic similarity to the known lethal RaTG13 virus.
- Eventually at some stage in December 2019, Wuhan Municipal Health Commission would notify the central CCP government in Beijing regarding the onset of the virus emergence among various patients. The approach by the Chinese government involved resorting to continuing the coverup of the outbreak, going to lengths to eliminate all remaining evidence and possible circulation of a suspected outbreak where possible.
- The Communist Party of China would then conspire with the World Health Organization in order to deter other member-states from investigating further into China and ultimately to assist with the concealment of the outbreak presence within the country. At this stage, efforts were being made to downplay the threat towards the international community.
- By January 2020, The Chinese People's Liberation Army Major General Chen Wei was appointed to oversee initial containment efforts over the city of Wuhan including supervision of the Wuhan Institute of Virology. Chen is also known as a leading military bio-warfare expert, which makes the motive for her selection likely due to the prospect that the city was under threat of biological agents. Several allegations made against WIV would immediately later be denied by the Chinese government.
- When the outbreak sufficiently circulated over multiple countries where it could be difficult to trace over a high number of suspected cases, it was deemed suitable for the World Health Organization to eventually inform the international community. The advise provided by WHO would however remain minimal to represent its true extent, encouraging governments to dismiss any pressing concerns over its spread. Given it was Chinese New Years (busiest travel periods to China) the Chinese government would also benefit from this traffic as having several cases spread across multiple countries would complicate investigations further.
- One of the largest information warfare campaigns would be launched in coordination with multiple parties ranging from governments, Big Tech and intergovernmental organisations to induce mass censorship. The aim would be to ensure the narrative could remain controlled to a proposed script, degrade contradicting hypothesises and efforts to amplify any connotation with China to be perceived as "xenophobic" or "fringe conspiracy" etc.
- At that point, the World Health Organisation biggest concern was over the ability of member-states to cooperate with it's own narrative and refrain against any dissent. Major efforts would then be made to help induce the removal of many heads of government that went against WHO. There is now a power-struggle and threat to state sovereignty between whether an intergovernmental body can detract loyalty when competing against it's own member-states.

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Sources:

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https://archive.ph/8QmFT
https://zenodo.org/record/4477081#.YIGAG-hKhPY
https://www.news-medical.net/news/20210217/The-origin-of-SARS-CoV-2-furin-cleavage-site-remains-a-mystery.aspx
https://www.dailymail.co.uk/news/article-9629563/Chinese-scientists-created-COVID-19-lab-tried-cover-tracks-new-study-claims.html
https://www.sciencealert.com/lab-or-nature-the-current-evidence-for-each-of-the-sars-cov-2-origin-theories
https://health.mountsinai.org/blog/did-sars-cov-2-emerge-from-nature-or-a-lab/
https://www.taiwannews.com.tw/en/news/4099020
https://jamanetwork.com/journals/jama/fullarticle/2762452
https://zslpublications.onlinelibrary.wiley.com/doi/abs/10.1111/j.1469-7998.1971.tb01323.x
https://www.nature.com/articles/nature12711
https://www.biorxiv.org/content/10.1101/2020.05.01.073262v1
https://www.preprints.org/manuscript/202102.0264/v1
https://www.sciencedirect.com/science/article/pii/S1873506120304165
https://www.nature.com/articles/s41586-020-2169-0
https://www.thelancet.com/journals/laninf/article/PIIS1473-3099(18)30006-9/fulltext
https://abcnews.go.com/Politics/trump-admin-pulls-nih-grant-coronavirus-research-ties/story?id=70418101
https://armscontrolcenter.org/wp-content/uploads/2016/02/Escaped-Viruses-final-2-17-14-copy.pdf
https://onlinelibrary.wiley.com/doi/full/10.1002/bies.202000091
https://www.franceculture.fr/sciences/covid-19-loms-percera-t-elle-le-mystere-de-lorigine-du-

virus
https://www.washingtonpost.com/politics/2021/05/25/timeline-how-wuhan-lab-leak-theory-suddenly-became-credible/
https://www.atlanticcouncil.org/wp-content/uploads/2021/02/Weaponized-How-rumors-about-COVID-19s-origins-led-to-a-narrative-arms-race.pdf#page=13
https://www.nature.com/articles/d41586-017-07766-9
https://www.nature.com/articles/nature.2017.21487
https://www.ncbi.nlm.nih.gov/books/NBK285579/
https://www.thelancet.com/action/showPdf?pii=S0140-6736%2820%2930183-5
https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7378494/

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